#16

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Atty. Docket: KIMCHI=2A

In re Application of:

Conf. No.: 4171

KIMCHI, et al

Art Unit: 1652

Appln. No.: 09/719,748

Examiner: M. Monshipouri

Filed: February 27, 2001

Washington, D.C.

For: DAP-KINASE RELATED PROTEIN)

April 10, 2003

DECLARATION UNDER 37 CFR \$1.132

Honorable Commissioner for Patents Washington, D.C. 20231

Sir:

I, Adi KIMCHI, hereby declare and state as follows:

I am the same Adi Kimchi who is the sole inventor of the invention disclosed and claimed in the above-identified application.

I am also the same Adi Kimchi who is listed among the co-authors of the publication, Boaz Inbal, Gidi Shani, Ofer Cohen, Joseph L. Kissil, and Adi Kimichi, Molecular and Cellular Biology 20:1044-1054 (2000), which publication is cited as the reference for SPTrEMBL database accession no. 075892 with a release date of November 8, 1998.

A copy of the printout from the SPTrEMBL database for accession no. 075892 is attached hereto. As can be seen from this printout, accession no. AAC35001 from the EMBL/Genbank

database, which is the same sequence as accession no. 075892, is cross-referenced. A printout of accession no. AAC35001 is attached hereto to show that it cites the same Inbal et al. reference as above and that it is the direct submission of A. Kimchi and B. Inbal on March 9, 1998.

While Boaz Inbal, Gidi Shani, Ofer Cohen, and Joseph 1. Kissil were co-authors with me on the publication and while Boaz Inbal was involved in the direct submission of the sequence, they were not involved in the conception of the invention and are not co-inventors of the invention claimed in the above-identified application no. 09/719,748.

made herein of my own knowledge are true and that all statements made on information and belief are believed to be true; and further that these statements were made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment, or both, under Section 1001 of Title 18 of the United States Code and that such willful false statements may jeopardize the validity of the application or any patent issued thereon.

14 5 03.

Adi Kinchi

NiceProt View of TrEMBL: O75892

General information about the entry

O75892 Entry name

O75892 Primary accession number

Secondary accession numbers

None

Entered in TrEMBL in Release 08, November 1998 Sequence was last modified in Release 08, November 1998

Annotations were last modified in Release 24, June 2003

Name and origin of the protein

DAP-kinase related protein 1 Protein name

None Synonyms Gene name None

Homo sapiens (Human) [TaxID: 9606] From

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Taxonomy

Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini;

Hominidae; Homo.

References

[1] SEQUENCE FROM NUCLEIC ACID.

TISSUE=Kidney;

MEDLINE=20094983; PubMed=10629061;

Inbal B., Shani G., Cohen O., Kissil J.L., Kimchi A.;

"Death-associated protein kinase-related protein 1, a novel Serine/Threonine kinase involved in apoptosis.";

Mol. Cell. Biol. 20:1044-1054(2000).

Comments

SIMILARITY: BELONGS TO THE SER/THR FAMILY OF PROTEIN KINASES.

Cross-references

EMBL	AF052941; AAC35001.1;
HSSP	Q63450; 1A06.
Genew	HGNC:2675; DAPK2.
CleanEx	HGNC:2675; DAPK2.
GO	GO:0005737; Cellular component: cytoplasmic chromosome (traceable author statement).

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T-965	P.008/008	

F-463

	GO:0005524; Molecular function: ATP binding activity (inferred from electronic annotation).				
	GO:0004683; Molecular function: calmodulin regulated protein kinase activity (traceable author statement).				
	GO:0004713; Molecular function: protein tyrosine kinase activity (inferred from electronic annotation).				
	GO:0016740; Molecular function: transferase activity (inferred from electronic annotation).				
	GO:0006917;Biological process: induction of apoptosis (traceable author statement).				
	GO:0006468; Biological process: protein amino acid phosphorylation (inferred from electronic annotation).				
InterPro	IPR000719; Prot_kinase. IPR002290; Ser_thr_pkinase. IPR001245; Tyr_pkinase.				
Pfam	PF00069; pkinase; 1.				
PRINTS	PR00109; TYRKINASE.				
ProDom	PD000001; Prot_kinase; 1. [Domain structure / List of seq. sharing at least 1 domain].				
SMART	SM00220; S_TKc; 1.				
PROSITE	PS00107; PROTEIN_KINASE_ATP; 1. PS50011; PROTEIN_KINASE_DOM; 1. PS00108; PROTEIN_KINASE_ST; 1.				
Implicit links to	Ensembl; ProtoMap; PRESAGE; ModBase; SWISS-2DPAGE.				

Keywords

ATP-binding; Kinase; Serine/threonine-protein kinase; Transferase.

Features

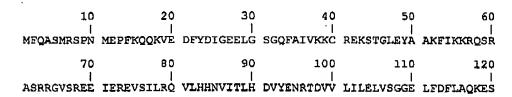
None

Sequence information

Length: 370 AA

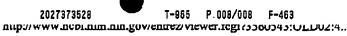
Molecular weight: 42923 Da

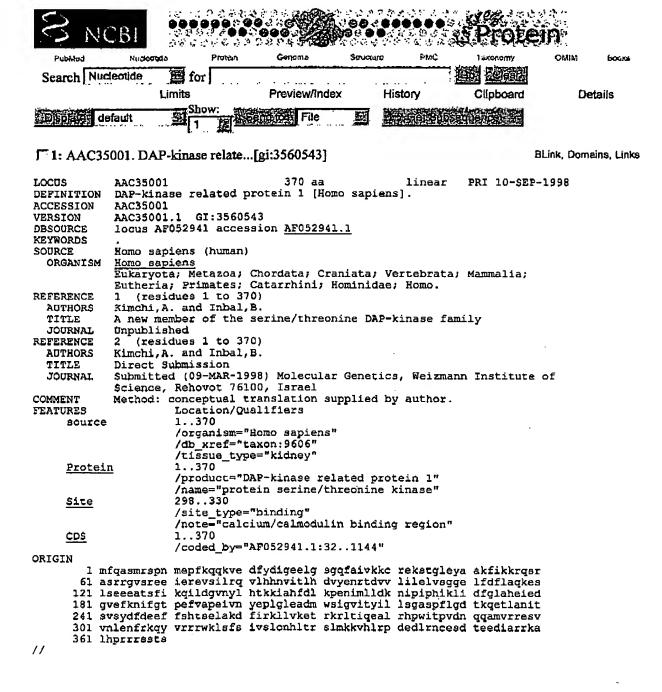
CRC64: 09502B4ADCD20F91 [This is a checksum on the sequence]



of 3

180 	170 	160 	150 	. 140	130
DFGLAHEIED	NIPIPHIKLI	KPENIMLLDK	HTKKIAHFDL	KOILDGVNYL	LSEEEATSFI
240 	230 I	220 	210	200	190 I
	LSGASPFLGD			PEFVAPEIVN	GVEFKNIFGT
300 <u>QQ</u> AMVRRESV	290 I RHPWITPVDN	280 RKRLTIQEAL	270 FIRKLLVKET	260 FSHTSELAKD	250 SVSYDFDEEF
360 TEEDIARRKA	350 DEDLRNCESD	340 I SLMKKVHLRP	330 IVSLCNHLTR	320 VRRRWKLSFS	310 VNLENFRKQY
					370 I LHPRRRSSTS





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